1/4

ATGGAGTCCGGGCTACTGCGGCCAGCGCCGGTGAGCGAGGTCATCGTCCTTCACTACAA CCGCAGTGTGCTTGGCTGTGCGCTTTCATCGTGCTGGAGAACCTGGCTGTGCTCTTG GTGCTGGGCCGCCATCCTCGCTTCCATGCACCCATGTTCCTGCTCCTGGGTAGTCTTAC CTTGTCGGACCTGCTCGCTGGGGCCGCCTACGCCACCAACATCCTGCTGTCCGGGCCGC TCACATTGCGCCTGTCGCCTGCGCTCTGGTTTGCGCGCGAAGGGGGTGTCTTCGTGGCG CTCGCAGCGTCGGTGCTGAGCCTCCTGGCCATTGCTCTAGAGCGCCACCTTACCATGGC CCGTCGTGGACCCGCGCCTGCCGCCAGTCGCGCTCGCACGCTGGCGATGGCGGTGGCCG CCTGGGGCTTGTCGCTGCTGCTGGGGCTGCCCGCGCTGGGCTGGAACTGCTTGGGA CGCCTGGAAGCCTGCTCCACCGTGCTGCCGCTCTACGCCAAGGCCTATGTGCTCTTCTG CGTGCTGGCCTTCCTGGGCATCCTGGCTGCCATCTGTGCGCTCTATGCAAGGATTTACT GTCAGGTGCGGGCCAACGCGCGTCGCCTGCGGGCCGGTCCTGGGTCCCGTAGGGCCACG TCCTCCTCGCGATCCCGGCACACGCCACGGTCGTTGGCCCTGCTCCGCACGCTTAGCGT GGTGCTCCTGGCCTTCGTGGCCTGCTGGGACCTCTATTTCTCTTGCTATTACTGGATG TCGCGTGCCCAGCCCGCGCGTGTCCTGTGCTTCTGCAAGCCGATCCCTTCCTGGGTCTA GCCATGGCTAACTCGCTGCTGAATCCTATCATCTACACCTTCACCAACCGAGACCTGCG CCACGCGCTCCTGAGGCTGCTCTGCTGTGGCCGCGGACCCTGCAACCAAGACTCCTCCA ACAGTTTGCAGCGATCCCCAAGTGCTGTTGGGCCTTCCGGTGGAGGCCTGCGACGCTGC CTGCCACCAACCCTGGATCGCAGCTCTAGCCCATCAGAACACTCGTGTCCCCAGCGGGA CGGGATGGACACCAGCTGCTCCACTGGCAGTCCCGGAGCAGCAACCGCCAACCGGACCC TGGTGCCTGATGCTACAGACTGA

FIG.1

MESGLLRPAPVSEVIVLHYNYTGKLRGARYQPGAGLRADAAVCLAVCAF
IVLENLAVLLVLGRHPRFHAPMFLLLGSLTLSDLLAGAAYATNILLSGP
LTLRLSPALWFAREGGVFVALAASVLSLLAIALERHLTMARRGPAPAAS
RARTLAMAVAAWGLSLLLGLLPALGWNCLGRLEACSTVLPLYAKAYVLF
CVLAFLGILAAICALYARIYCQVRANARRLRAGPGSRRATSSSRSRHTP
RSLALLRTLSVVLLAFVACWGPLFLLLLDVACPARACPVLLQADPFLG
LAMANSLLNPIIYTFTNRDLRHALLRLLCCGRGPCNQDSSNSLQRSPSA
VGPSGGGLRRCLPPTLDRSSSPSEHSCPQRDGMDTSCSTGSPGAATANR
TLVPDATD

FIG.2

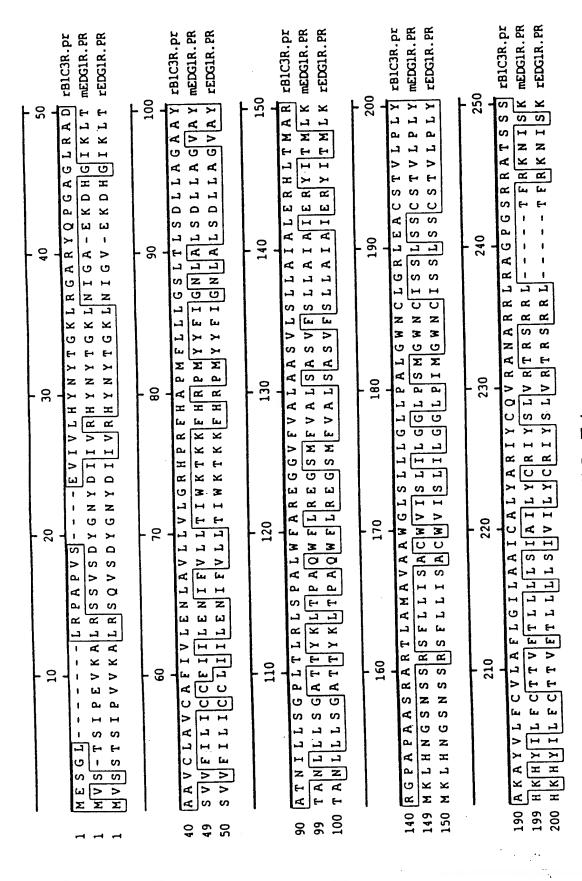


FIG.3A

rB1C3R.pr mEDG1R.PR rEDG1R.PR	rBlC3R.pr mEDG1R.PR rEDG1R.PR	rBlC3R.pr mEDG1R.PR rEDG1R.PR	rBlC3R.pr mEDG1R.PR rEDG1R.PR
SVVLLAFVACWGPLFLLLLLDVACPARACPVLLQA TBIC3R.pr IIVLSVFIACWAPLFILLLLDVGCKAKTCDILYKA MEDGIR.PR IIVLSVFIACWAPLFILLLDVGCKAKTCDILYKA MEDGIR.PR	320 350 350 350 350 350 350 X T F T N R D L R H A L L R L L C C G R G P C N Q D S S N S L Q R R D C R L L L L C C C R G P C N Q D S S N S L Q R R L L L R I V S C C K - C P N R EDGIR.PR Y T L T N K E M R R A F I R I I S C C K - C P N r EDGIR.PR	PSEHSC-PORDGMDTSCSTGSPGAAKSDNSSHPOKDDGDNPETIMSSGNVKSDDGDNPETIMSSGNV	다 본 다
12 > >	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	( ) ( ) ( )	
240 RSRHTPRSLALLRT 244 GSRSSEKSLALLKT 245 ASRSSEKSLALLKT	310 290 DPFLGLAMANSLLN 294 EYFLVLAVLNSGTN 295 EYFLVLAVLNSGTN	340 S P S A V G P S G G L R R 334 G D S A G K F K R 335 G D S A G K F K R	410 389 TANRTLVPDATD 379NSSS 380NSSS

FIG.3B